# Jianle Sun

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## **Education**

## **Shanghai Jiao Tong University**

Sep. 2021-Jun. 2024 (expected)

- Master of Science (Research-based) in Biostatistics & Bioinformatics
- Advisors: Associate Prof. Yue Zhang & Prof. Zhangsheng Yu
- **GPA:** major 3.83/4.0, total 3.90/4.0
- **Related courses:** Fundamental Mathematical Statistics (A+), Biology Mathematics I (A+), Applied Stochastic Processes (A+), Causal Inference Methods in Data Science (A+), Data Mining (A+), Introduction to Computational Biology (A+), Machine Learning (A), Statistical Learning (A), Medical Bioinformatics (A), Neural Network Theory and Application (A-)

### **Shanghai Jiao Tong University**

Sep. 2017- Jun. 2021

- Bachelor of Science in Biotechnology (Bioinformatics)
- **GPA** 3.94/4.3 (rank 1/25)
- Related courses:
  - Quantitative: Calculus I & II (A-), Linear Algebra (A+), Probability & Statistics (A), Thinking and Approach of Programming (A), Biostatistical Models (A+), Biostatistics and Mathematical Modeling (A), Stochastic Simulation Methods and Its Applications (A), Business Analytics and Data Mining (A)
  - **Biology-related:** Inorganic and Analytic Chemistry, Organic Chemistry, Physical Chemistry, Biochemistry I & II, Genetics, Microbiology, Anatomy and Physiology, Ecology and Evolution, Cell Biology, Molecular Biology, Immunology, Developmental Biology
- Thesis: Bayesian Network-based Mendelian Randomization for Causal Inference (A)

### **Research Interest**

Causal Inference & Causal Learning, Statistical Genetics and Genomics, Statistical & Computational Methods in Multi-omics, Genetic Epidemiology, Bayesian Methods

# **Publications & Working Papers**

- **Sun, J.**, Lyu, R., Deng, L., Li, Q., Zhao, Y., & Zhang, Y. (2022). SMetABF: A rapid algorithm for Bayesian GWAS meta-analysis with a large number of studies included. *PLOS Computational Biology*, *18*(3), e1009948. https://doi.org/10.1371/journal.pcbi.1009948
- Sun, J., Zhou, J., Gong, Y., Pang, C., Ma, Y., Zhao, J., Yu, Z., & Zhang, Y. Bayesian network-based Mendelian randomization for variant prioritization and phenotypic causal inference. *Under Review*
- Sun, J., Deng, L., Li, Q., Zhou, J., & Zhang, Y. Dynamic relations between longitudinal morphological, behavioral, and emotional indicators and cognitive impairment: evidence from the Chinese Longitudinal Healthy Longevity Survey. *Under Review*
- Lyu, R., **Sun, J.**, Xu, D., Jiang, Q., Wei, C., & Zhang, Y. (2021). GESLM algorithm for detecting causal SNPs in GWAS with multiple phenotypes. *Briefings in Bioinformatics*, 22(6), bbab276. <a href="https://doi.org/10.1093/bib/bbab276">https://doi.org/10.1093/bib/bbab276</a>
- · Zhou, Y., Fa, B., Wei, T., **Sun, J.**, Yu, Z., & Zhang, Y. (2021). Elastic Correlation Adjusted Regression (ECAR) scores for high dimensional variable importance measuring. *Scientific Reports*, *II*(1), 1-12. <a href="https://doi.org/10.1038/s41598-021-02706-0">https://doi.org/10.1038/s41598-021-02706-0</a>
- · Gong, Y., Xu, J., Gao, R., **Sun, J.**, Yu, Z., & Zhang, Y. Single-Cell Biclustering for Cell-Specific Transcriptomic Perturbation Detection in AD Progression. *Under Review* <a href="https://dx.doi.org/10.2139/ssrn.4493250">https://dx.doi.org/10.2139/ssrn.4493250</a> (preprint)
- · Liang, C., Bai, W., Qiao, L., Ren, Y., **Sun, J.**, Ye, P., Yan, H., Ma, X., Zuo, W., & Ouyang, W. Rethinking the BERT-like pretraining for DNA sequences. *ICLR in submission*

# **Projects**

- Bayesian GWAS meta-analysis methodology and its application to Parkinson's disease

- · Improve the subset selection process in ABF-based Bayesian GWAS meta-analysis by introducing MCMC and Shotgun Stochastic Search algorithm
- · Analyzing genetic patterns of Parkinson's disease and other autoimmune disorders through cross-trait Bayesian meta-analysis
- Implementing proposed algorithms as an R package and an R shiny online software

### Bayesian network-based Mendelian randomization for causal inference

- · Proposing the random graph forest, comprised of a series of Bayesian network structure learning processes, to prioritize candidate variants and select proper genetic instruments, and obtained pleiotropy-robust causal estimates through Bayesian Mendelian randomization with a shrinkage prior
- Appling proposed model in UK Biobank to infer the causality between erythrocyte-related hematologic parameters and blood pressures, and between lymphocyte and psychiatric disorders like schizophrenia

# Epidemiological analysis on cognitive impairment using Chinese Longitudinal Healthy Longevity Survey (CLHLS) datasets

- · Identifying risk factors of aging cognitive impairment with data of participants' physical indicators, activities of daily life, emotional status, and cognitive abilities from 8 waves of CLHLS datasets
- Establishing dynamic predictive models for cognitive impairment using Bayesian joint model and dynamic-DeepHit, a deep learning-based survival model
- Assessing longitudinal indicators' effect on cognitive decline by controlling confounders using marginal structural Cox model.

# Dynamic Mendelian randomization with genomics and single-cell transcriptomics

- Extending Mendelian randomization to single-cell resolution by integrating eQTL and single-cell transcriptomics
- Employing functional data analysis and g-estimation of longitudinal confounding to model dynamic effect in single-cell trajectory

# Deep learning-based prospective stroke predictions using dynamic longitudinal physiological records

- · Labeling longitudinal physiological records according to symptom descriptions in each follow-up
- Employing Cox model with lasso to select crucial risk factors, RNN module to capture the longitudinal features, and random forest for the final classification on stroke

## Internship

## Shanghai AI laboratory | Research Intern on AI for Science

Jul. 2023 – Oct. 2023

- Developing pretrained BERT-based DNA large language model
- · Conducting downstream tasks, including identifying genomic functional elements (promoters, enhancers, etc.), predicting epigenomic & expressive features, and assessing the effects of variants

# · Baiyi Health | Biodata Analyst

Aug. 2022 - Jan. 2023

- Establishing biomarker-based biological age models for eight systems/organs, and establishing classification model for thyroid disorders
- · Participating in the trial design and data analysis for the performance of Eye Parameter Evaluator

#### Honors

- · Graduation with honor: College Graduate Excellence Award of Shanghai, 2021 (5%)
- · National Encouragement Scholarship
- Fan Hsu-chi Scholarship
- · Tang You Shuqi Named Scholarship
- · Sino Biological Named Scholarship
- Gold award and nominated for best new application project in the International Genetically Engineered Machine (iGEM) Competition, 2019

# **Skills**

- Language: Chinese, English (IELTS 7.5), Japanese (Junior), learning experience in ancient languages like Latin
- Computer: Python, R, MATLAB, SPSS, LaTeX, PyTorch, Linux, PLINK